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Location/Qualifiers  
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/mol\_type="unassigned DNA"  
/db\_xref="taxon:9606"

ORIGIN

Query Match 100.0%; Score 20; DB 6; Length 467;  
Best Local Similarity 100.0%; Pred. No. 21;  
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Y 1 ATCGCATGATATCGCATGAT 20  
420 ATCGCATGATATCGCATGAT 401

RESULT 2  
AX364976/c  
LOCUS AX364976 Sequence 127 from Parent MO0206315. 468 bp DNA linear PAT 15-FEB-2002  
DEFINITION AX364976  
ACCESSION AX364976  
VERSION AX364976.1 GI:18696866  
KEYWORDS  
SOURCE Homo sapiens (human)  
ORGANISM Homo sapiens  
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.  
REFERENCE 1 Mirtz, L., Freilich, S. and Bernstein, J.  
TITLE Novel nucleic acid and amino acid sequences  
JOURNAL Patent: WO 0206315-A 127 24-JAN-2002;  
Compugen Ltd. (IL)  
LOCATION/Qualifiers  
1..468  
/organism="Homo sapiens"  
/mol\_type="unassigned DNA"  
/db\_xref="taxon:9606"

ORIGIN

Query Match 100.0%; Score 20; DB 6; Length 468;  
Best Local Similarity 100.0%; Pred. No. 21;  
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Y 1 ATCGCATGATATCGCATGAT 20  
421 ATCGCATGATATCGCATGAT 402

RESULT 3  
S82024/c  
LOCUS S82024 696 bp mRNA linear PRI 03-AUG-1996  
DEFINITION SCG10=neuron-specific growth-associated protein/stathmin homolog  
[human, embryo, mRNA, 696 nt].  
S82024  
S82024.1 GI:1478502  
ACCESSION S82024  
VERSION S82024  
KEYWORDS  
SOURCE Homo sapiens (human)  
ORGANISM Homo sapiens  
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
Mammalia; Eutheria; Primates; Catarrhini; Homnidae; Homo.  
REFERENCE 1 (bases 1 to 696)  
AUTHORS Okazaki, T., Wang, H., Masliah, E., Cao, M., Johnson, S.A., Sundsmo, M.,  
Saitoh, T. and Mori, N.  
TITLE SCG10, a neuron-specific growth-associated protein in Alzheimer's  
disease.  
JOURNAL Neurobiol. Aging 16 (6): 883-894 (1995)  
MEDLINE 96192979  
PUBMED 8622778  
REMARK GenBank staff at the National Library of Medicine created this  
entry [NCBI gibbed 177683] from the original journal article.  
This sequence comes from Fig. 1.  
Location/Qualifiers

source  
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/organism="Homo sapiens"  
/mol\_type="mRNA"  
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gene  
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CDS  
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homolog. This sequence comes from Fig. 1"

ORIGIN

Query Match 100.0%; Score 20; DB 9; Length 696;  
Best Local Similarity 100.0%; Pred. No. 20;  
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Y 1 ATCGCATGATATCGCATGAT 20  
649 ATCGCATGATATCGCATGAT 630

RESULT 4  
AY006733/c  
LOCUS AY006733 376 bp DNA linear BCT 19-SEP-2000  
DEFINITION Uncultured rumen bacterium 16S ribosomal RNA gene, partial  
sequence.  
AY006733  
AY006733.1 GI:10189345  
ACCESSION AY006733  
VERSION AY006733  
KEYWORDS  
SOURCE uncultured rumen bacterium  
ORGANISM uncultured rumen bacterium  
Bacteria; environmental samples.  
REFERENCE 1 (bases 1 to 376)  
AUTHORS Tamalis, D., Dyer, D., Ralph, D., Hartman, K., Phillips, W., Coleman, S.  
TITLE Assessing diversity in bovine rumen microflora in response to  
feeding using 16S ribosomal RNA sequencing  
JOURNAL Unpublished  
2 (bases 1 to 376)  
AUTHORS Tamalis, D., Dyer, D., Ralph, D., Hartman, K., Phillips, W., Coleman, S.  
TITLE Direct Submission  
JOURNAL Submitted (07-AUG-2000) Microbiology and Immunology, Oklahoma  
University Health Sciences Center, EMSB 1053, PO Box 26901,  
Oklahoma City, OK 73190, USA  
LOCATION/Qualifiers  
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/organism="uncultured rumen bacterium"  
/mol\_type="genomic DNA"  
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/db\_xref="taxon:136703"  
/note="from winter wheat forage-fed steer #169"  
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/product="16S ribosomal RNA"

ORIGIN

Query Match 84.0%; Score 16.8; DB 1; Length 376;  
Best Local Similarity 90.0%; Pred. No. 9.5e+02;  
Matches 18; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Y 1 ATCGCATGATATCGCATGAT 20  
187 AGCGCAGATATCGCATGAT 168

CompuGen Ltd. (IL)  
Location/Qualifiers  
source 1..467  
/organism="Homo sapiens"  
/mol\_type="unassigned DNA"  
/db\_xref="taxon:9606"

## ORIGIN

Query Match 100.0%; Score 20; DB 6; Length 467;  
Best Local Similarity 100.0%; Pred. No. 21;  
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Y 1 ATCATGGCATATCATGGCAT 20  
DB 401 ATCATGGCATATCATGGCAT 420

## RESULT 2

AX364976 468 bp DNA linear PAT 15-FEB-2002  
LOCUS Sequence 127 from Patent WO0206315.  
ACCESSION AX364976  
VERSION AX364976.1 GI:18696866  
KEYWORDS

SOURCE Homo sapiens (human)  
ORGANISM Homo sapiens

REFERENCE Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
Mammalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.  
1

AUTHORS Mintz, L., Freilich, S. and Bernstein, J.  
TITLE Novel nucleic acid and amino acid sequences  
JOURNAL Patent: WO 0206315-A 127 24-JAN-2002;  
CompuGen Ltd. (IL)  
Location/Qualifiers

FEATURES  
source 1..468  
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/mol\_type="unassigned DNA"  
/db\_xref="taxon:9606"

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Query Match 100.0%; Score 20; DB 6; Length 468;  
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Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Y 1 ATCATGGCATATCATGGCAT 20  
DB 402 ATCATGGCATATCATGGCAT 421

## RESULT 3

S82024 696 bp mRNA linear PRI 03-AUG-1996  
LOCUS SCG10-neuron-specific growth-associated protein/stathmin homolog  
DEFINITION [human, embryo, mRNA, 696 nt].  
ACCESSION S82024  
VERSION S82024.1 GI:1478502  
KEYWORDS

SOURCE Homo sapiens (human)  
ORGANISM Homo sapiens

REFERENCE Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
Mammalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.  
1 (bases 1 to 696)  
AUTHORS Okazaki, T., Wang, H., Masliak, E., Cao, M., Johnson, S.A., Sundemo, M.,  
Saitoh, T. and Mori, N.

## TITLE

SCG10, a neuron-specific growth-associated protein in Alzheimer's  
disease

JOURNAL Neurobiol. Aging 16 (6), 883-894 (1995)  
MEDLINE 96192979  
PubMed 8622778

## REMARK

Genbank staff at the National Library of Medicine created this  
entry [NCBI gisdbeg 177683] from the original journal article.  
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## source

1..696  
/organism="Homo sapiens"  
/mol\_type="mRNA"  
/db\_xref="taxon:9606"  
1..696  
/gene="SCG10"  
29..568  
/gene="SCG10"

## CDS

/note="neuron-specific growth-associated protein/stathmin  
homolog; this sequence comes from Fig. 1"

/product="SCG10"  
/protein\_id="AA86428.1"  
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ERHAAEVRNKEQLVELSG"

## ORIGIN

Query Match 100.0%; Score 20; DB 9; Length 696;  
Best Local Similarity 100.0%; Pred. No. 20;  
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Y 1 ATCATGGCATATCATGGCAT 20  
DB 630 ATCATGGCATATCATGGCAT 649

## RESULT 4

AY006733 376 bp DNA linear BCT 19-SEP-2000  
LOCUS AY006733  
DEFINITION Uncultured rumen bacterium 16S ribosomal RNA gene, partial  
sequence.  
ACCESSION AY006733  
VERSION AY006733.1 GI:10189345  
KEYWORDS

## SOURCE

ORGANISM uncultured rumen bacterium  
uncultured rumen bacterium  
Bacteria; environmental samples.

REFERENCE 1 (bases 1 to 376)  
TAMAILIS, D., DYER, D., RALPH, D., HARTMAN, K., PHILLIPS, W., COLEMAN, S.  
and Iandolo, J.  
TITLE Assessing diversity in bovine rumen microflora in response to  
feeding using 16S ribosomal RNA sequencing  
JOURNAL Unpublished  
2 (bases 1 to 376)

## AUTHORS

TAMAILIS, D., DYER, D., RALPH, D., HARTMAN, K., PHILLIPS, W., COLEMAN, S.  
and Iandolo, J.  
TITLE Direct Submission  
JOURNAL Submitted (07-AUG-2000) Microbiology and Immunology, Oklahoma  
University Health Sciences Center, BMSB 1053, PO Box 26901,  
Oklahoma City, OK 73190, USA  
Location/Qualifiers

## FEATURES

source 1..376  
/organism="uncultured rumen bacterium"  
/mol\_type="genomic DNA"  
/specific\_host="Bos taurus"  
/db\_xref="taxon:136703"  
/note="from winter wheat forage-fed steer #169"  
1..376  
/product="16S ribosomal RNA"

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Best Local Similarity 90.0%; Pred. No. 9.5e+02;  
Matches 18; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Y 1 ATCATGGCATATCATGGCAT 20  
DB 168 ATCATGGCATATCATGGCAT 167